

TECHNICAL AMENDMENTS TO THE SPECIFICATION:

Please amend Tables 1 and 2 on pp. 51 and 52, respectively, of the instant specification as indicated hereinbelow:

Table 1. Sequence of gene specific primers used in the C-ORF cloning technology.

Primer	Gene	SEQ ID NO	Sequence
AP-1	D-SLAP	1	5'-TTCTGGTCGACTAGTGTTTAAACTCGAGAC-3' (SEQ ID NO:1)
AP-2	D-SLAP	2	5'-CACGATCAGTCGACGAAGTCGACACTCGAG-3' (SEQ ID NO:2)
M5R1	mda-5	3	5'-TTTTTTTTTTTTCAGAGTAAAACAATC-3' (SEQ ID NO:3)
M5R2	mda-5	4	5'-TGTGCACCATCATTGTTCCCAAGCC-3' (SEQ ID NO:4)
M5R3	mda-5	5	5'-AATCTGACATTGGACTCATTGAC-3' (SEQ ID NO:5)
M5R4	mda-5	6	5'-GTTTGAATCCTTGTCATTATTTCTAG-3' (SEQ ID NO:6)
M5S1	mda-5	7	5'-GCCTGAGAGCCCTGTGGACAACCTCG-3' (SEQ ID NO:7)
56R1	ISG-56	8	5'-GTGGCTGATATCTGGGTGCCTAAGG-3' (SEQ ID NO:8)
56R2	ISG-56	9	5'-CCTAAGGACCTTGTCACAGAGTTC-3' (SEQ ID NO:9)
56S1	ISG-56	10	5'-CCAGATCTCAGAGGAGCCTGGCTAAGC-3' (SEQ ID NO:10)
M9R1	mda-9	11	5'-AATCAGGATAAAGTGTCAACTATC-3' (SEQ ID NO:11)
M9R2	mda-9	12	5'-ATCCCAAAGTAGCTAGGTACATAATC-3' (SEQ ID NO:12)
M9S1	mda-9	13	5'-CCTCAGAAGTCCGTGCCAGTGACCGG-3' (SEQ ID NO:13)
FnR1	fibronectin	14	5'-TTTTTTTTTTTGTGGAATGTAAATC-3' (SEQ ID NO:14)
FnR2	fibronectin	15	5'-AGATGGATCTTGGCAGAGAGACATGC-3' (SEQ ID NO:15)
PCTAR1	PCTA-1	16	5'-GAAGAAGTAGAACATCAGTGCC-3' (SEQ ID NO:16)
PCTAR2	PCTA-1	17	5'-TCTTCTGTACAGCAGTATCTTACAT-3' (SEQ ID NO:17)
PCTAR3	PCTA-1	18	5'-TTTTTTTTTTTTTTTTTGTGTTGCATGCGG-3' (SEQ ID NO:18)
PCTAR4	PCTA-1	19	5'-TTACAAACAGCTCCCAAATGGTGAAACT-3' (SEQ ID NO:19)

Table 2. Sequence of the 5' end of the C-ORF products.

Name	SEQ ID NO	5' sequence	G/C score	% G/C
mda-20	20	5'-GCGCGCCGGCCT-3' (SEQ ID NO:20)	11/12	91.7
ISG-56	21	5'-TGCAGAACGGCT-3' (SEQ ID NO:21)	7/12	58.3
mda-9	22	5'-GGCGGCGGCGGC-3' (SEQ ID NO:22)	12/12	100.0
PCTA-1A	23	5'-TGGAGGCCTGGA-3' (SEQ ID NO:23)	8/12	66.7
PCTA-1B	24	5'-GCCAGTGCCTCA-3' (SEQ ID NO:24)	8/12	66.7
PCTA-1C	25	5'-CGATGTGGCCTT-3' (SEQ ID NO:25)	7/12	58.3
OLD-35	26	5'-CGGAGGACCAAT-3' (SEQ ID NO:26)	7/12	58.3
PSGen 12	27	5'-GCGGTGGTGACG-3' (SEQ ID NO:27)	9/12	75.0
PSGen 28	28	5'-GTGTGGTGTGTC-3' (SEQ ID NO:28)	7/12	58.3
PSGen 42	29	5'-GGCGTTGCGACG-3' (SEQ ID NO:29)	9/12	75.0
G/C/ score	--	8994867986 74	85/120	70.8